



Figure 2-1

AccI	243	GT'mk_AC (0 Err) - 2 Fragment(s)
		909
ApaI	402	G_GGCC'C (0 Err) - 2 Fragment(s)
		750
BamHI	200	G'GATC_C (0 Err) - 2 Fragment(s)
		952
EcoRI	343	G'AATT_C (0 Err) - 2 Fragment(s)
		809
HindIII	135	A'AGCT_T (0 Err) - 2 Fragment(s)
		1017
NciI	196	CC's_GG (0 Err) - 2 Fragment(s)
		956
NcoI	60	C'CATG_G (0 Err) - 4 Fragment(s)
		135 288 669
PstI	337	C_TGCA'G (0 Err) - 3 Fragment(s)
		366 449
PvuII	570	CAG'CTG (0 Err) - 2 Fragment(s)
		582
Sali	242	G'TCGA_C (0 Err) - 2 Fragment(s)
		910

== Linear Map of Sequence:

```

1  ggggcaggcaatgagagctgcactctggctggggaaggcatgagtgacagacccacagca 60
   ccccgctccgttactctcgacgtgagaccgaccccttcgctactcactgtctcgggtgtcgt
   ^      *      ^      *      ^      *      ^      *      ^      *
61  aggcgggtggggtaagtgtggacctttgtgtaccagagagaacatcatgggtggctttcaaa 120
   tccgccacccattcacacctggaaacacatgggtctctcttgtagtaccaccgaaagttt
   ^      *      ^      *      ^      *      ^      *      ^      *
                                HindIII
                                /
121 ggggtctggactcaagctttctgaaagcagtcacagcggaatttctggccatgcttatt 180
   ccccgacctgagttcgaaagacctttcgtcagtggtcgccttaagaccggtacgaataa
   ^      *      ^      *      ^      *      ^      *      ^      *
                                BamHI
                                /
181 tttgttctcctcagcctgggatccaccatcaactggggtggaacagaaaagcctttaccg 240
   aaacaagaggagtcggacctaggtggtagttgacccaccttgtcttttcggaatggc
   ^      *      ^      *      ^      *      ^      *      ^      *
Sali

```


Figure 2-3

```

841 aaagctgccagcaaaacaaaggaagctacatggaggtggaggacaacaggagtcaggta 900
    ttcgacgggtcgtttgtttcccttcgatgtacctccacctcctgtgtcctcagtcct
        ^      *      ^      *      ^      *      ^      *      ^      *
                                           NciI
                                           \
901 gagacggatgacctgattctaaaacctggagtggtgcattgattgacgttgaccgggga 960
    ctctgcctactggactaagattttggacctcaccacgtacactaactgcaactggccct
        ^      *      ^      *      ^      *      ^      *      ^      *
                                           ^      *
961 gaggagaagaaggggaaagaccaatctggagaggtattgtcttcagtatgactagaagat 1020
    ctctcttcttcccccttctgggttagacctctccataacagaagtcatactgatcttcta
        ^      *      ^      *      ^      *      ^      *      ^      *
                                           ^      *
1021 cgactgaaagcagacaagactccttagaactgtcctcagatttccttccaccattaag 1080
    gcgtgactttcgtctgttctgaggaatcttgacaggagtctaaaggaagtggttaattc
        ^      *      ^      *      ^      *      ^      *      ^      *
                                           ^      *
1081 gaaacagatttggttataaattagaaatgtgcaggtttggtgtttcatgtcatattactca 1140
    ctttgctaaacaatatattaatctttacacgtccaaacaacaaagtacagtataatgagt
        ^      *      ^      *      ^      *      ^      *      ^      *
                                           ^      *
1141 gtctaaacaata (SEQ ID NO:1) 1200
    cagatttggttat
        ^      *      ^      *      ^      *      ^      *      ^      *
  
```

===== End of Analysis =====

Return	Help	Report Bugs	
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Citation

Algorithm Citation:

Not given.

Program Citation:

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